

Expression cloning of cDNA encoding a human β -1,3-*N*-acetylglucosaminyltransferase that is essential for poly-*N*-acetylactosamine synthesis

Katsutoshi Sasaki*, Kazumi Kurata-Miura*, Minoru Ujita†, Kiyohiko Angata†, Satoshi Nakagawa*, Susumu Sekine*, Tatsunari Nishi*, and Minoru Fukuda†,‡

*Tokyo Research Laboratories, Kyowa Hakko Kogyo Co., Ltd., Machida, Tokyo 194, Japan; and †Glycobiology Program, The Burnham Institute, La Jolla Cancer Research Center, La Jolla, CA 92037

Edited by Helen M. Ranney, Alliance Pharmaceutical Corp., San Diego, CA, and approved October 27, 1997 (received for review August 27, 1997)

- ♦ ABSTRACT
- ♦ INTRODUCTION
- ♦ EXPERIMENTAL PROCEDURES
- ♦ RESULTS
- ♦ DISCUSSION
- ♦ FOOTNOTES
- ♦ ACKNOWLEDGEMENTS
- ♦ REFERENCES

- ▶ Abstract of this Article
- ▶ Reprint (PDF) Version of this Article
- ▶ Similar articles found in:
PNAS Online
ISI Web of Science
PubMed
- ▶ PubMed Citation
- ▶ This Article has been cited by:
other online articles
- ▶ Search Medline for articles by:
Sasaki, K., || Fukuda, M.
- ▶ Search for citing articles in:
ISI Web of Science (18)
- ▶ Alert me when:
new articles cite this article
- ▶ Download to Citation Manager

ABSTRACT

The structure and biosynthesis of poly-*N*-acetylactosamine display a dramatic change during development and oncogenesis. Poly-*N*-acetylactosamines are also modified by various carbohydrate residues, forming functional oligosaccharides such as sialyl Le^x. Herein we describe the isolation and functional expression of cDNA encoding β -1,3-*N*-acetylglucosaminyltransferase (iGnT), an enzyme that is essential for the formation of poly-*N*-acetylactosamine. For this expression cloning, Burkitt lymphoma Namalwa KJM-1 cells were transfected with cDNA libraries derived from human melanoma and colon carcinoma cells. Transfected Namalwa cells overexpressing the i antigen were continuously selected by fluorescence-activated cell sorting because introduced plasmids containing Epstein-Barr virus replication origin can be continuously amplified as episomes. Sibling selection of plasmids recovered after the third consecutive sorting resulted in a cDNA clone that directs the increased expression of i antigen on the cell surface. The deduced amino acid sequence indicates that this protein has a type II membrane protein topology found in almost all mammalian glycosyltransferases cloned to date. iGnT, however, differs in having the longest transmembrane domain among glycosyltransferases cloned so far. The iGnT transcripts highly expressed in fetal brain and kidney and adult brain but expressed ubiquitously in various adult tissues. The expression of the presumed catalytic domain as a fusion protein with the IgG binding domain of protein A enabled us to demonstrate that the cDNA encodes iGnT, the enzyme responsible for the formation of $\text{GlcNAc}\beta 1 \rightarrow 3\text{Gal}\beta 1 \rightarrow 4\text{GlcNAc} \rightarrow \text{R}$ structure and poly-*N*-acetylactosamine extension.

INTRODUCTION

Poly-*N*-acetylactosamine is a unique carbohydrate composed of *N*-acetylactosamine repeats ($\text{Gal}\beta 1 \rightarrow 4\text{GlcNAc}\beta 1 \rightarrow 3$)_n. Poly-*N*-acetylactosamines can be attached to *N*-glycans, *O*-glycans, or glycolipids and provide the backbone structure for additional modifications, which are very often cell-type-specific oligosaccharide structures (1–3). In humans, fetal erythrocytes express a linear poly-*N*-acetylactosamine, $\text{Gal}\beta 1 \rightarrow 4\text{GlcNAc}\beta 1 \rightarrow 3\text{Gal}\beta 1 \rightarrow 4\text{GlcNAc} \rightarrow \text{R}$, and adult erythrocytes express a branched poly-*N*-acetylactosamine, $\text{Gal}\beta 1 \rightarrow 4\text{GlcNAc}\beta 1 \rightarrow 3(\text{Gal}\beta 1 \rightarrow 4\text{GlcNAc}\beta 1 \rightarrow 6)\text{Gal}\beta 1 \rightarrow 4\text{GlcNAc} \rightarrow \text{R}$ (4, 5). These poly-*N*-acetylactosamines are the determinants for the i and I antigen, which are the first human alloantigens shown to display developmental change (6, 7). For the synthesis of

poly-*N*-acetylglucosamine, β -1,3-galactosylglucosaminyltransferase (iGnT) is essential for the formation of the i antigen (Fig. 1). When I-forming β -1,6-*N*-acetylglucosaminyltransferase (IGnT) is also present with iGnT, the I antigen is synthesized instead of the i antigen (4, 8–14).

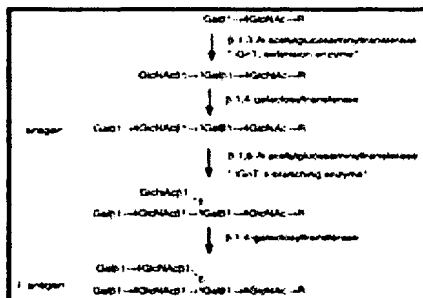


Fig. 1. Structure and biosynthesis of i and I antigens. The i antigen is synthesized by iGnT followed by β -1,4-galactosyltransferase. The i antigen is converted to the I antigen by the stepwise addition of a GlcNAc β 1 \rightarrow 6 and a Gal β 1 \rightarrow 4 residue (8). As an alternative pathway, another IGnT adds β -1,6-*N*-acetylglucosamine to GlcNAc β 1 \rightarrow 3Gal β 1 \rightarrow 4GlcNAc β 1 \rightarrow 6R precursor and this product, GlcNAc β 1 \rightarrow 3(GlcNAc β 1 \rightarrow 6)Gal β 1 \rightarrow 4GlcNAc β 1 \rightarrow 6R, is converted to Gal β 1 \rightarrow 4GlcNAc β 1 \rightarrow 3(Gal β 1 \rightarrow 4GlcNAc β 1 \rightarrow 6)Gal β 1 \rightarrow 4GlcNAc β 1 \rightarrow 6R, forming I antigen (9) (adapted from refs. 4 and 8–12).

[View Larger Version of this Image (OK GIF file)]

In granulocytes, monocytes, and memory T lymphocytes, poly-*N*-acetylglucosamines carry sialyl Le^x, NeuNAc α 2 \rightarrow 3Gal β 1 \rightarrow 4(Fuc α 1 \rightarrow 3)GlcNAc β 1 \rightarrow 6R, at their termini (15), which plays a critical role in recruiting leukocytes to inflammatory sites (16–19) and possibly when tumor cells adhere at metastatic sites (20–22). As an antigen specific to mouse embryo, stage specific embryonic antigen (SSEA)-1 antigen was identified as Gal β 1 \rightarrow 4(Fuc α 1 \rightarrow 3)GlcNAc β 1 \rightarrow 6R, which is present in poly-*N*-acetylglucosamines. By using anti-SSEA-1 antibody or oligosaccharides containing Gal β 1 \rightarrow 4(Fuc α 1 \rightarrow 3)GlcNAc terminus as inhibitors, it was demonstrated that SSEA-1 might participate in adhesive events that are involved in compaction during embryogenesis (23, 24).

By expression cloning, we have isolated a cDNA encoding IGnT by using Chinese hamster ovary (CHO) cells containing polyoma large tumor antigen because CHO cells lack the I antigen (12). In contrast, it has been extremely difficult to clone cDNA encoding iGnT because this enzyme is ubiquitously present in various cells although its abundance is not as high as β -1,4-galactosyltransferase (10, 11, 13, 14).

Recently, we have demonstrated that it is possible to clone GD3/GT3 synthase by using COS-1/GD3 cells by transient expression cloning (25). COS-1/GD3 cells synthesized a substantial amount of GD3 but only a small amount of GT3. A cDNA encoding GD3/GT3 synthase could be isolated from COS-1/GD3 cells that were overexpressing GD3 and GT3 after transfection (25). This cloning taught us that it is possible to clone a cDNA encoding iGnT as long as transfected cells become highly positive for iGnT. This is important because cells entirely lacking the i antigen were not available. For this expression cloning strategy, we have selected Namalwa KJM-1 cells that express Epstein-Barr virus nuclear antigen 1 (EBNA-1) (26–28). In cells expressing EBNA-1, plasmids containing the replication origin of Epstein-Barr virus oriP can be continuously amplified as episomes (29), thus allowing continuous selection of those cells overexpressing the i antigen. Herein we report this cloning strategy, resulting in the isolation of a cDNA encoding iGnT, which is essential for poly-*N*-acetylglucosamine synthesis.

EXPERIMENTAL PROCEDURES

Isolation of a Human iGnT cDNA Clone.

All of the various cell lines in our hands, including COS-1 and CHO cells, were found to be positive for the i antigen. We thus decided to clone iGnT by overexpressing iGnT in cells where the i antigen was present. For this, Namalwa KJM-1, a human Burkitt lymphoma cell line, was used as recipient cells. cDNA libraries derived from poly(A)⁺ RNA of human melanoma WM266-4 (26) and colonic carcinoma SW1116 (27) cell lines were constructed in pAMo vector containing oriP of the Epstein-Barr virus. Namalwa KJM-1 cells were then transfected with a mixture of the above cDNA libraries and the transfected cells were first selected in the presence of G418, because pAMo also contains a neomycin-resistance gene. After 13 days, the transfected cells were incubated with human anti-i antigen serum (Den; ref. 7) followed by fluorescein isothiocyanate-conjugated goat anti-human IgM. The i-antigen-positive cells were isolated by fluorescence-activated cell sorting (EPICS Elite flow cytometer, Coulter) and cultured for an additional 18 days. The transfected cells were sorted two more times by using the same procedure.

After the third sorting, plasmids were recovered by the Hirt procedure (30) from those cells that were highly positive for the i antigen expression. Among 18 clones of 75 plasmids recovered, clone, 16-2-12, was found to increase the expression of the i antigen by a factor of 7 compared with the cells transfected with pAMo containing no cDNA insert. The cDNA in this plasmid was sequenced by the dideoxynucleotide chain-termination method (31). The cDNA insert in 16-2-12 was digested with *Hind*III and *Xba*I and cloned into the *Hind*III and *Eco*RV sites of pcDNA3.1 (Invitrogen), resulting in pcDNA3.1-iGnT.

Northern Blot Analysis of Various Human Tissues.

Human multiple tissue Northern blots of poly(A)⁺ RNA were purchased from CLONTECH, and these blots were hybridized with a gel-purified cDNA insert of pcDNA3.1-iGnT or pcDNAI-iGnT (2) after labeling with [α -³²P]dCTP by random-oligonucleotide priming (Prime-IT II labeling kit, Stratagene).

Construction and Expression of the Protein A-iGnT Fusion Vector.

The cDNA fragment encoding the stem region plus putative catalytic domain of iGnT was prepared by PCR using pcDNA3.1-iGnT as a template and fused with cDNA encoding a signal peptide sequence and the IgG binding domain of *Staphylococcus aureus* protein A (25–28). 5' and 3' primers for this PCR are 5'-CGGATCCACGGTCCGTGGACCAGGT-3' and 5'-TCGCTCGAGGGCTCAGCAGCGTCGGG-3' (*Bam*HI and *Xho*I sites are underlined). The PCR product encoding amino acid residues 53–415 of iGnT was ligated into the *Bam*HI and *Xho*I sites of pcDNAI-A (32), yielding plasmid pcDNAI-A?iGnTc. Plasmid pcDNAI-A and pcDNAI-A?iGnTc were separately transfected with LipofectAmine (GIBCO/BRL) into COS-1 cells as described (32), and 48 hr after transfection the medium was replaced with serum-free medium, macrophage-SFM (GIBCO/BRL) and cultured for an additional 24 hr. The chimeric iGnTc secreted into the culture medium was adsorbed to IgG-Sepharos 6FF (Pharmacia) and the enzyme bound to the beads was used as an enzyme source (33). Alternatively, the culture medium was concentrated by 10-fold and directly used as an enzyme source.

iGnT Assays and Product Characterization.

iGnT assays were performed essentially as described (34) except for a few modifications. In all assays, the reaction mixtures contained 0.1 mM UDP-[³H]GlcNAc (1×10^6 cpm/nmol), 20 mM MnCl₂, 5 mM ATP, and 5 mM acceptor oligosaccharide, lacto-*N*-neotetraose, in a final volume of 100 μ l of 100 mM cacodylate buffer (pH 7.0). In addition, 10 mM *N*-acetylglucosamine-1,5-lactone was included to inhibit breakdown of products by β -hexosaminidase(s) when the culture medium was used as an enzyme source. After 10-hr incubation at 37° C, 0.3 ml of QAE-Sephadex was added to the reaction mixture. The supernatant recovered was again mixed with QAE-Sephadex, and its derived supernatant was applied to a column (1.0 \times 120 cm) of Bio-Gel P-4 equilibrated with 0.1 M NH₄HCO₃. The above product, purified after Bio-Gel P-4 gel filtration, was incubated with β -1,4-galactosyltransferase (5 milliunits Boehringer Mannheim) in the same reaction mixture as described above, except that UDP-[³H]GlcNAc was replaced with 0.5 mM UDP-[³H]Gal (1×10^6 cpm/5 nmol). α ₁-Acid glycoprotein (100 μ g), desialylated by neuraminidase treatment, was also used as an acceptor. To analyze its product, α ₁-acid glycoprotein was precipitated by adding ethanol (90% final concentration) to the reaction mixture and the centrifuged and the precipitate after dissolving in 0.1 M NH₄HCO₃ was applied to a column (1.0 \times 27 cm) of Sephadex G-50 (superfine). These biosynthetic products were digested with jack bean β -*N*-acetylglucosaminidase, β -galactosidase, or endo- β -galactosidase (35) and subjected to Bio-Gel P-2 gel filtration equilibrated with 0.1 M NH₄HCO₃.

RESULTS

Isolation of a cDNA Clone that Determines the Expression of the i Antigen.

Our preliminary results indicated that there is no cell line available that is deficient in poly-*N*-acetyllactosamine synthesis judging from the staining with human anti-i serum (Den; ref 7) or tomato lectin, which reacts with poly-*N*-acetyllactosamine (36). By using tomato lectin, we also attempted to isolate CHO cells that are defective in iGnT without success. We thus decided to use Namalwa KJM-1 cells that express EBNA-1 as recipient cells for expression cloning. Namalwa KJM-1 cells have been used for cloning of α -2,3-sialyltransferase (ST3Gal IV), Fuc-TVII, and GD3 synthase (26–28). Besides, plasmids containing oriP continuously replicate as episomes in the presence of EBNA-1. This latter advantage allowed us to enrich Namalwa KJM-1 cells expressing an increased amount of iGnT without isolation of plasmids from sorted cells and retransfection of those plasmids into the recipient cells.

As a source of mRNAs for construction of a cDNA library, we tested HL-60, WM266-4, and SW1116 cells, which were known or shown to express the i antigen. As shown in Fig. 2, the expression of the i antigen on transfected Namalwa KJM-1 cells was substantially increased when the cells were transfected with a mixture of cDNA libraries derived from WM266-4 and SW1116. In contrast, no increase of the i antigen expression was achieved by using HL-60 cDNA library (data not shown). After the third sorting, more than 50% of the transfected cells showed increased expression of the i antigen (Fig. 2a). Plasmids were isolated from these Namalwa KJM-1 cells that were highly positive for the i antigen, and then each clone was tested for the ability to increase the expression of the antigen. One of the plasmids, 16-2-12, directed 7 times-increased expression of the i antigen, and an unrelated plasmid, 16-2-3 did not (Fig. 2b). Because Namalwa KJM-1 cells endogeneously express antigen, i antigen was detected in mock-transfected cells (Fig. 2b).

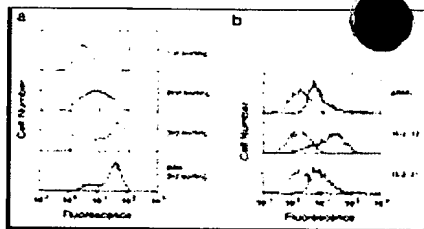


Fig. 2. Expression cloning of *i* enzyme by flow cytometry. *(a)* Namalwa KJM-1 cells stably transfected with the WM266-4 and SW1116 cDNA libraries were stained with the anti-*i* antibody (Den) and subjected to three rounds of sorting with a fluorescence-activated cell sorter. The cells with strong fluorescence intensities indicated by the bars were collected and subjected to the subsequent sorting. *(b)* Flow cytometric analysis of the cells transfected with the empty vector pAMo (*Top*), a plasmid 16-2-12 harboring *iGnT* cDNA (*Middle*), or an unrelated plasmid 16-2-31 (*Bottom*) after staining with the

anti-*i* antibody (solid lines) or PBS (dotted lines).

[View Larger Version of this Image (OK GIF file)]

Predicted Amino Acid Sequence of *iGnT*.

The cDNA insert in plasmid 16-2-12 encoding *iGnT* contains an ORF predicting a protein of 415 amino acid residues (Fig. 3). A hydropathy plot predicts that this protein has a type II transmembrane topology, with a short cytoplasmic sequence at the amino terminus, followed by the transmembrane domain, and then by the so-called stem region and a large catalytic domain, which presumably resides in the Golgi lumen. This topology has been found in almost all mammalian glycosyltransferases so far cloned [17]. The *iGnT* is, however, characteristic in having a relatively long transmembrane domain that likely spans from residue 9 to residue 36, consisting of 28 residues. It is also characteristic in having only one basic amino acid at each end (Arg at residue 8 and His at residue 37) that flanks the transmembrane domain.

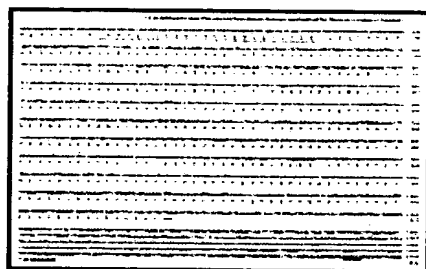


Fig. 3. DNA and translated amino acid sequences of *iGnT*. The full-length nucleotide and amino acid sequences of *iGnT* are shown. The signal/membrane-anchoring domain is underlined. Potential N-glycosylation sites are marked by asterisks. A polyadenylation signal is doubly underlined. The sequences are numbered relative to the translation initiation site.

[View Larger Version of this Image (OK GIF file)]

There are two potential *N*-glycosylation sites (Fig. 3, asterisks). A consensus sequence for polyadenylation signal is present at nucleotides 1902-1907, which is probably used, judging from the size of mRNA (see below). No significant similarity was found between this protein sequence and other sequences reported in the Gene Data Bank.

Expression of *iGnT* and *IGnT* mRNAs in Human Tissues.

To determine the expression profile of *iGnT* and *IGnT*, Northern blots of poly(A)⁺ RNA derived from various human tissues were hybridized with *iGnT* probe followed by *IGnT* probe [12]. A band of 2.2 kb for *iGnT* transcript was detected in all poly(A)⁺ RNA isolated from various tissues (Fig. 4). In fetal tissues, the signal was more prominent for poly(A)⁺ RNA derived from brain and kidney than from lung or liver. In adult tissues, the signal was weaker from thymus, peripheral blood leukocytes, lung, and liver than the other tissues.

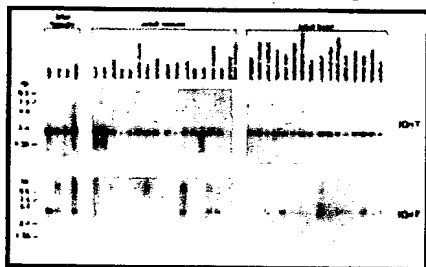


Fig. 4. Northern blot analysis of *iGnT* and *IGnT* in various human fetal and adult tissues. Each lane contained 2 μ g of poly(A)⁺ RNA. The same blots were probed by ³²P-labeled *iGnT* cDNA (*iGnT*) followed by *IGnT* cDNA (*IGnT*).

[View Larger Version of this Image (OK GIF file)]

In general, however, all of the tissues examined contained the transcript for *iGnT*. In contrast, the signal for *IGnT* was found only in certain tissues (Fig. 4). In fetal tissues, the transcript for *IGnT* was substantially expressed in brain and moderately expressed in kidney and lung but was almost undetectable in liver. In adult tissue, the transcript for *IGnT* was strongest in prostate, moderate in small intestine and colon, and barely detected in heart, brain, kidney, and pancreas. In adult brain, the *IGnT* transcript is much more prominent in cerebellum than the other parts of brain. Thus, these results indicate that *iGnT* is ubiquitously expressed in various tissues, although its amount varies according to tissues, and *IGnT* is clearly expressed in a tissue-specific manner.

Expression of Catalytically Active iGnT.

To confirm that the cDNA isolated encodes iGnT, the sequence corresponding to the putative stem region plus catalytic domain, iGnTc, was fused in-frame with cDNA encoding a signal peptide and the IgG binding domain of *S. aureus* protein A (25–28). As shown in Fig. 5A, the spent medium derived from COS-1 cells transfected with pcDNA1-A?iGnTc produced [3 H]GlcNAc-labeled α_1 -acid glycoprotein. Less than one-fifth of the product was obtained by using the supernatant from COS-1 cells transfected with pcDNA1-A lacking cDNA insert. The labeled product was digested by endo- β -galactosidase, resulting in the release of [3 H]GlcNAc $\beta_1 \rightarrow$ 3Gal (Fig. 5C). The activity in the supernatant from mock-transfected COS-1 cells was most likely due to iGnT endogenously expressed in COS-1 cells (38). In parallel, the labeled α_1 -acid glycoprotein was incubated with UDP-[3 H]Gal and β -1,4-galactosyltransferase, and the resultant product was digested by endo- β -galactosidase, releasing [3 H]Gal $\beta_1 \rightarrow$ 4[3 H]GlcNAc $\beta_1 \rightarrow$ 3Gal (Fig. 5C). The structures of these released oligosaccharides were deduced based on the standard oligosaccharide and the substrate specificity of endo- β -galactosidase (35). To confirm that the above activity was derived from the expressed enzyme, the chimeric enzyme adsorbed to IgG-Sepharose was used as an enzyme source. Fig. 5B demonstrates that the same labeled product was obtained by this enzyme but the IgG-Sepharose beads adsorbed by the mock-transfected culture medium, had no activity.

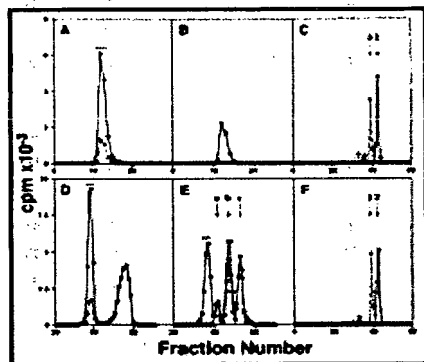


Fig. 5. Gel filtration analysis of reaction products by iGnT. COS-1 cells were transfected with pcDNA1-A?iGnT (solid line) or pcDNA1-A (dotted line), and the derived spent medium (A and D) or the enzyme adsorbed to IgG-Sepharose (B) was used as an enzyme source. (A and B) The desialylated α_1 -acid glycoprotein was used as an acceptor and the product was analyzed by Sephadex G-50 gel filtration. (C) Bio-Gel P-2 gel filtration after endo- β -galactosidase digestion of product A obtained in A (solid line) or the galactosylated product A (dotted line). (D) Lacto-*N*-neotetraose was used as an acceptor and the product was analyzed by Bio-Gel P-4 gel filtration. The product shown in D (denoted by horizontal bar) was incubated with β -1,4-galactosyltransferase and UDP-[3 H]Gal and subjected to Bio-Gel P-4 gel filtration. Peaks a, b, and c denote contaminated radioactivity derived from UDP-[3 H]galactose present in both experiments with (solid line) or without (dotted line) the acceptor. (F) Bio-Gel P-2 gel filtration after endo- β -galactosidase treatment of the product obtained in D (solid line) or the product obtained in E (dotted line). Arrows 2 and 3 in C and F denote the elution positions of GlcNAc $\beta_1 \rightarrow$ 3Gal and Gal $\beta_1 \rightarrow$ 4GlcNAc $\beta_1 \rightarrow$ 3Gal, respectively. The products subjected to additional analysis are shown by horizontal bars. [View Larger Version of this Image (OK GIF file)]

To determine whether iGnT adds GlcNAc to *N*-acetylglucosamine repeats, lacto-*N*-neotetraose was used as an acceptor. As shown in Fig. 5D, iGnT efficiently added GlcNAc to lacto-*N*-neotetraose (0.7 nmol formed from 500 nmol of the acceptor) and this product [3 H]GlcNAc $\beta_1 \rightarrow$ 3Gal $\beta_1 \rightarrow$ 4GlcNAc $\beta_1 \rightarrow$ 3Gal $\beta_1 \rightarrow$ 4Glc was converted to [3 H]Gal $\beta_1 \rightarrow$ 4[3 H]GlcNAc $\beta_1 \rightarrow$ 3Gal $\beta_1 \rightarrow$ 4Glc by incubation with β -1,4-galactosyltransferase and UDP-[3 H]Gal (Fig. 5E). The same results were obtained with the chimeric enzyme adsorbed to IgG-Sepharose beads (data not shown). From these labeled products, [3 H]GlcNAc $\beta_1 \rightarrow$ 3Gal and [3 H]Gal $\beta_1 \rightarrow$ 4[3 H]GlcNAc $\beta_1 \rightarrow$ 3Gal were respectively released by endo- β -galactosidase treatments (Fig. 5F).

In addition, the chimeric enzyme adsorbed to IgG-Sepharose transferred β -1,3-linked GlcNAc to Gal $\beta_1 \rightarrow$ 4Glc, but the IgG-Sepharose beads, adsorbed by the mock-transfected culture medium, had no activity. No transfer of [3 H]GlcNAc was observed toward Gal $\beta_1 \rightarrow$ 3GlcNAc $\beta_1 \rightarrow$ 3Gal $\beta_1 \rightarrow$ 4Glc (data not shown). Thus, these results indicate that iGnT cloned in the present study can add *N*-acetylglucosamine to lactose and both *N*-acetylglucosamines attached to *N*-glycans of α_1 -acid glycoprotein and lacto-*N*-neotetraose.

DISCUSSION

In the present study, we report the isolation of a human cDNA clone encoding iGnT, determination of its expression in various tissues, and demonstration of its *in vitro* activity. For this cloning, we used Namalwa KJM-1 cell line as recipient cells for transfection and a vector containing the replication origin of Epstein-Barr virus oriP. Because Namalwa KJM-1 cells synthesize EBNA-1, amplification of a vector containing oriP is possible in an episomal manner. Because of this advantage, it was possible to continuously enrich the transfected cells expressing an increased amount of the i antigen without isolation of plasmids and reintroduction of the plasmids to recipient cells. This is an improved method and combination of two previous methods for cloning GD3/GT3 synthase and Fuc-TVII by using an overexpression protocol (25, 27) and cloning ST3Gal IV and GD3 synthase (26, 28) by using Namalwa KJM-1 cells and pAMo vector.

The predicted amino acid sequence of iGnT has several characteristics. In particular, iGnT apparently has a longer transmembrane domain consisting of 28 amino acid residues than other glycosyltransferases so far cloned. Most other glycosyltransferases contain a transmembrane domain consisting of 14–24 amino acid residues (17). As shown previously, the size of poly-*N*-acetylglucosamines is longer in those attached to membrane glycoproteins than those attached to secretory glycoproteins (1). Moreover, human chorionic gonadotropin α glycoprotein acquired poly-*N*-acetylglucosamine once it became a membrane glycoprotein by fusing with the transmembrane and cytoplasmic portions of vesicular stomatitis virus G protein (18). These results strongly suggest that iGnT has a unique characteristic in binding to acceptor substrates, preferentially adding poly-*N*-acetylglucosamine to membrane glycoproteins. Such a unique property may be related to its long transmembrane domain. The availability of iGnT cDNA will allow us to determine whether or not changes in its transmembrane domain alter the property of iGnT in this aspect.

In previous studies, iGnT was detected and partially purified from human serum (10), Novikoff ascites tumor cells (13), and calf serum (14). The substrate specificity of these enzymes are very similar to those of the cloned iGnT, and all the enzymes add *N*-acetylglucosamine to $\text{Ga}\beta 1 \rightarrow 4\text{GlcNAc}$, $\text{Ga}\beta 1 \rightarrow 4\text{Glc}$, and $\text{Ga}\beta 1 \rightarrow 4\text{GlcNAc}\beta 1 \rightarrow 3\text{Ga}\beta 1 \rightarrow 4\text{Glc}$ but not to $\text{Ga}\beta 1 \rightarrow 3\text{GlcNAc}\beta 1 \rightarrow 3\text{Ga}\beta 1 \rightarrow 4\text{Glc}$. These results strongly suggest that the enzyme cloned in the present study can initiate the synthesis of poly-*N*-acetylglucosamine and also elongate poly-*N*-acetylglucosamine. On the other hand, the iGnT cloned in the present study has a predicted molecular weight of 47,125 for its polypeptide, whereas the molecular mass of iGnT partially purified from calf serum was found to be 70 kDa (14). If two *N*-glycans are attached to the cloned iGnT, then the mature iGnT may have an approximate mass of 53 kDa. These results suggest that the calf serum iGnT may be different from the human iGnT cloned in the present study. Alternatively, iGnT may contain a large amount of *O*-glycans in addition to *N*-glycans, although this possibility is unlikely judging from the deduced amino acid sequence.

It has been shown that tumor cells express more poly-*N*-acetylglucosamine than normal counterparts (19–42). It has been also demonstrated that the activity of GnTV is also increased in those tumor cells (39–41). In parallel, iGnT was shown to prefer $\text{Ga}\beta 1 \rightarrow 4\text{GlcNAc}\beta 1 \rightarrow 6\text{Man}\alpha 1 \rightarrow 6\text{Man}\beta 1 \rightarrow \text{R}$, which is formed by GnTV, as an acceptor over other side chains in *N*-glycans (13, 43, 44). These combined results suggest that the tumorigenic phenotype acquired by the introduction of GnTV cDNA (45) could be due to the increase of poly-*N*-acetylglucosamine built on the side chain synthesized by GnTV. Further studies using the cloned iGnT will be of significance to determine whether the increase of poly-*N*-acetylglucosamine, but not the increase in the side chain formed by GnTV, is the primary cause for transformed phenotype displayed by tumor cells.

FOOTNOTES

† To whom reprint requests should be addressed at: The Burnham Institute, 10901 North Torrey Pines Road, La Jolla, CA 92037. e-mail: minoru@licrf.edu

This paper was submitted directly (Track II) to the *Proceedings* Office.

Abbreviations: iGnT, β -1,3-*N*-acetylglucosaminyltransferase; IGnT, I-branch forming β -1,6-*N*-acetylglucosaminyltransferase; R, aglycon; EBNA-1, the nuclear antigen-1 of Epstein-Barr virus; GnTV, *N*-acetylglucosaminyltransferase ∇ .

Data deposition: The sequence reported in this paper has been deposited in the GenBank database (accession no. AF029893).

ACKNOWLEDGEMENTS

We thank Dr. Michiko N. Fukuda for useful discussion, Ms. Sachiko Kodama and Ayumi Natsume for excellent technical assistance, Dr. Edgar Ong for critical reading of the manuscript, and Ms. Susan Greaney for organizing the manuscript. The work was supported by Grant CA48737 from the National Cancer Institute (to M.F.) and a Toyobo Biotechnology Fellowship (to M.U.).

REFERENCES

1. Fukuda, M. & Fukuda, M. N. (1994) in *The Biology of Glycoproteins*, ed. Ivatt J. (Plenum, New York), pp. 183–234.
2. Feizi, T. (1985) *Nature (London)* **314**, 53–57 [ISI][Medline].
3. Hakomori, S. (1989) *Adv. Cancer Res.* **52**, 257–331 [ISI][Medline].
4. Fukuda, M., Fukuda, M. N. & Hakomori, S. (1979) *J. Biol. Chem.* **254**, 3700–3703 [ISI][Medline].
5. Watanabe, K., Hakomori, S., Childs, R. A. & Feizi, T. (1979) *J. Biol. Chem.* **254**, 3221–3228 [ISI][Medline].
6. Wiener, A. S., Unger, L. H., Cohen, L. & Feldman, J. (1956) *Ann. Intern. Med.* **44**, 221–240.
7. Feizi, T., Childs, R. A., Watanabe, K. & Hakomori, S.-I. (1979) *J. Exp. Med.* **149**, 975–980 [ISI][Abstract].
8. Gu, J., Nishikawa, A., Fujii, S., Gasa, S. & Taniguchi, N. (1992) *J. Biol. Chem.* **267**, 2994–2999 [ISI][Abstract].
9. Piller, F., Cartron, J. P., Maranduba, A., Veyrieres, A., Leroy, Y. & Fournet, B. (1984) *J. Biol. Chem.* **259**, 13385–13390 [ISI][Abstract].
10. Piller, F. & Cartron, J. P. (1983) *J. Biol. Chem.* **258**, 12293–12299 [ISI][Abstract].
11. Koenderman, A. H., Koppen, P. L. & Van den Eijnden, D. H. (1987) *Eur. J. Biochem.* **166**, 199–208 [ISI][Abstract].
12. Bierhuizen, M. F. A., Mattei, M. G. & Fukuda, M. (1993) *Genes Dev.* **7**, 468–478 [ISI][Abstract].
13. Van den Eijnden, D. H., Koenderman, A. H. & Schiphorst, W. E. (1988) *J. Biol. Chem.* **263**, 12461–12471 [Abstract].
14. Kawashima, H., Yamamoto, K., Osawa, T. & Irimura, T. (1993) *J. Biol. Chem.* **268**, 27118–27126 [ISI][Abstract].
15. Fukuda, M., Spooncer, E., Oates, J. E., Dell, A. & Klock, J. C. (1984) *J. Biol. Chem.* **259**, 10925–10935 [ISI][Abstract].
16. Lowe, J. B., Stoolman, L. M., Nair, R. P., Larsen, R. D., Berhend, T. L. & Marks, R. M. (1990) *Cell* **63**, 475–484 [ISI][Medline].
17. Phillips, M. L., Nudelman, E., Gaeta, F. C. A., Perez, M., Singhai, A. K., Hakomori, S.-I. & Paulson, J. C. (1990) *Science* **250**, 1130–1132 [ISI][Medline].
18. Walz, G., Aruffo, A., Kolanus, W., Bevilacqua, M. & Seed, B. (1990) *Science* **250**, 1132–1135 [ISI][Medline].
19. Lowe, B. J. (1994) in *Molecular Glycobiology*, eds. Fukuda, M. & Hindsgaul, O. (Oxford Univ. Press, Oxford), pp. 163–205.
20. Fukushima, K., Hirota, M., Terasaki, P. I., Wakisaka, A., Togashi, H., Chia, D., Suyama, N., Fukushi, Y., Nudelman, E. & Hakomori, S. (1984) *Cancer Res.* **44**, 5279–5285 [ISI][Medline].
21. Nakamori, S., Kameyama, M., Imaoka, S., Furukawa, H., Ishikawa, O., Sasaki, Y., Kabuto, T., Iwanaga, T., Matsushita, Y. & Irimura, T. (1993) *Cancer Res.* **53**, 3632–3637 [ISI][Medline].
22. Sawada, R., Tsuboi, S. & Fukuda, M. (1994) *J. Biol. Chem.* **269**, 1425–1431 [ISI][Abstract].
23. Bird, J. M. & Kimber, S. J. (1984) *Dev. Biol.* **104**, 449–460 [ISI][Medline].
24. Fenderson, B. A., Zehavi, U. & Hakomori, S. (1984) *J. Exp. Med.* **160**, 1591–1596 [ISI][Abstract].
25. Nakayama, J., Fukuda, M. N., Hirabayashi, Y., Kanamori, A., Sasaki, K., Nishi, T. & Fukuda, M. (1996) *J. Biol. Chem.* **271**, 3684–3691 [ISI][Abstract/Full Text].
26. Sasaki, K., Watanabe, E., Kawashima, K., Sekine, S., Dohi, T., Oshima, M., Hanai, N., Nishi, T. & Hasegawa, M. (1993) *J. Biol. Chem.* **268**, 22782–22787 [ISI][Abstract].
27. Sasaki, K., Kurata, K., Funayama, K., Nagata, M., Watanabe, E., Ohta, S., Hanai, N. & Nishi, T. (1994) *J. Biol. Chem.* **269**, 14730–14737 [ISI][Abstract].
28. Sasaki, K., Kurata, K., Kojima, N., Kurosawa, N., Ohta, S., Hanai, N., Tsuji, S. & Nishi, T. (1994) *J. Biol. Chem.* **269**, 15950–15956 [ISI][Abstract].
29. Margolskee, R. F., Kavathas, P. & Berg, P. (1988) *Mol. Cell. Biol.* **8**, 2837–2847 [ISI][Medline].
30. Hirt, B. (1967) *J. Mol. Biol.* **26**, 365–369 [Medline].
31. Sanger, F., Nicklen, S. & Coulson, A. R. (1977) *Proc. Natl. Acad. Sci. USA* **74**, 5463–5467 [ISI][Medline].
32. Nakayama, J. & Fukuda, M. (1996) *J. Biol. Chem.* **271**, 1829–1832 [ISI][Abstract/Full Text].
33. Bierhuizen, M. F. A. & Fukuda, M. (1992) *Proc. Natl. Acad. Sci. USA* **89**, 9326–9330 [ISI][Abstract].
34. Lee, N., Wang, W.-C. & Fukuda, M. (1990) *J. Biol. Chem.* **265**, 20476–20487 [ISI][Abstract].
35. Fukuda, M. N. (1981) *J. Biol. Chem.* **256**, 3900–3905 [ISI][Abstract].
36. Merkle, R. K. & Cummings, R. D. (1987) *J. Biol. Chem.* **262**, 8179–8189 [ISI][Abstract].
37. Schachter, H. (1994) in *Molecular Glycobiology*, eds. Fukuda, M. & Hindsgaul, O. (IRL Press, Oxford), pp. 88–162.
38. Fukuda, M., Guan, J.-L. & Rose, J. K. (1988) *J. Biol. Chem.* **263**, 5314–5318 [ISI][Abstract].
39. Yamashita, K., Ohkura, T., Tachibana, Y., Takasaki, S. & Kobata, A. (1984) *J. Biol. Chem.* **259**, 10834–10840 [ISI][Abstract].
40. Pierce, M. & Arango, J. (1986) *J. Biol. Chem.* **261**, 10772–10777 [ISI][Abstract].
41. Dennis, J. W., Lafert, S., Waghorn, C., Breitman, M. L. & Kerbel, R. S. (1987) *Science* **236**, 582–585 [ISI][Medline].
42. Saitoh, O., Wang, W.-C., Lotan, R. & Fukuda, M. (1992) *J. Biol. Chem.* **267**, 5700–5711 [ISI][Abstract].
43. Cummings, R. D. & Kornfeld, S. (1984) *J. Biol. Chem.* **259**, 6253–6260 [ISI][Abstract].
44. Sasaki, H., Bothner, B., Dell, A. & Fukuda, M. (1987) *J. Biol. Chem.* **262**, 12059–12076 [ISI][Abstract].
45. Demetriou, M., Nabi, I. R., Coppolino, M., Dedhar, S. & Dennis, J. W. (1995) *J. Cell. Biol.* **130**, 383–392 [ISI][Abstract].

Copyright (C)1997 by The National Academy of Sciences of the USA.
0027-8424/97/9414294-6\$2.00/0

This article has been cited by other articles:

- Ohyama, C., Smith, P. L., Angata, K., Fukuda, M. N., Lowe, J. B., Fukuda, M. (1998). Molecular Cloning and Expression of GDP-D-mannose-4,6-dehydratase, a Key Enzyme for Fucose Metabolism Defective in Lec13 Cells. *J. Biol. Chem.* 273: 14582-14587 [Abstract] [Full Text]
- Mattila, P., Salminen, H., Hirvas, L., Niittymäki, J., Salo, H., Niemelä, R., Fukuda, M., Renkonen, O., Renkonen, R. (1998). The Centrally Acting beta 1,6N-Acetylglucosaminyltransferase (GlcNAc to Gal). FUNCTIONAL EXPRESSION, PURIFICATION, AND ACCEPTOR SPECIFICITY OF A HUMAN ENZYME INVOLVED IN MIDCHAIN BRANCHING OF LINEAR POLY-N-ACETYLACTOSAMINES. *J. Biol. Chem.* 273: 27633-27639 [Abstract] [Full Text]
- Ujita, M., McAuliffe, J., Schwientek, T., Almeida, R., Hindsgaul, O., Clausen, H., Fukuda, M. (1998). Synthesis of Poly-N-acetylactosamine in Core 2 Branched O-Glycans. THE REQUIREMENT OF NOVEL beta -1,4-GALACTOSYLTRANSFERASE IV AND beta -1,3-N-ACETYLGLUCOSAMINYLTRANSFERASE. *J. Biol. Chem.* 273: 34843-34849 [Abstract] [Full Text]
- Ujita, M., McAuliffe, J., Suzuki, M., Hindsgaul, O., Clausen, H., Fukuda, M. N., Fukuda, M. (1999). Regulation of I-Branched Poly-N-Acetylactosamine Synthesis. CONCERTED ACTIONS BY i-EXTENSION ENZYME, I-BRANCHING ENZYME, AND beta 1,4-GALACTOSYLTRANSFERASE I. *J. Biol. Chem.* 274: 9296-9304 [Abstract] [Full Text]
- Ujita, M., McAuliffe, J., Hindsgaul, O., Sasaki, K., Fukuda, M. N., Fukuda, M. (1999). Poly-N-acetylactosamine Synthesis in Branched N-Glycans Is Controlled by Complemental Branch Specificity of i-Extension Enzyme and beta 1,4-Galactosyltransferase I. *J. Biol. Chem.* 274: 16717-16726 [Abstract] [Full Text]
- Schwientek, T., Nomoto, M., Levery, S. B., Merkx, G., van Kessel, A. G., Bennett, E. P., Hollingsworth, M. A., Clausen, H. (1999). Control of O-Glycan Branch Formation. MOLECULAR CLONING OF HUMAN cDNA ENCODING A NOVEL beta 1,6-N-ACETYLGLUCOSAMINYLTRANSFERASE FORMING CORE 2 AND CORE 4. *J. Biol. Chem.* 274: 4504-4512 [Abstract] [Full Text]
- Schwientek, T., Yeh, J.-C., Levery, S. B., Keck, B., Merkx, G., van Kessel, A. G., Fukuda, M., Clausen, H. (2000). Control of O-Glycan Branch Formation. MOLECULAR CLONING AND CHARACTERIZATION OF A NOVEL THYMUS-ASSOCIATED CORE 2 beta 1,6-N-ACETYLGLUCOSAMINYLTRANSFERASE. *J. Biol. Chem.* 275: 11106-11113 [Abstract] [Full Text]
- Ujita, M., Misra, A. K., McAuliffe, J., Hindsgaul, O., Fukuda, M. (2000). Poly-N-acetylactosamine Extension in N-Glycans and Core 2- and Core 4-branched O-Glycans Is Differentially Controlled by i-Extension Enzyme and Different Members of the beta 1,4-Galactosyltransferase Gene Family. *J. Biol. Chem.* 275: 15868-15875 [Abstract] [Full Text]
- Steffensen, R., Carlier, K., Wiels, J., Levery, S. B., Stroud, M., Cedergren, B., Nilsson Sojka, B., Bennett, E. P., Jersild, C., Clausen, H. (2000). Cloning and Expression of the Histo-blood Group Pk UDP-galactose:Galbeta 1-4Glcbeta 1-Cer alpha 1,4-Galactosyltransferase. MOLECULAR GENETIC BASIS OF THE p PHENOTYPE. *J. Biol. Chem.* 275: 16723-16729 [Abstract] [Full Text]
- Nakayama, J., Yeh, J.-C., Misra, A. K., Ito, S., Katsuyama, T., Fukuda, M. (1999). Expression cloning of a human alpha 1,4-N-acetylglucosaminyltransferase that forms GlcNAc alpha 1right-arrow4Galbeta right-arrowR, a glycan specifically expressed in the gastric gland mucous cell-type mucin. *Proc. Natl. Acad. Sci. U. S. A.* 96: 8991-8996 [Abstract] [Full Text]
- Zhou, D., Dinter, A., Gallego, R. G., Kamerling, J. P., Vliegthart, J. F. G., Berger, E. G., Hennot, T. (1999). A beta -1,3-N-acetylglucosaminyltransferase with poly-N-acetylactosamine synthase activity is structurally related to beta -1,3-galactosyltransferases. *Proc. Natl. Acad. Sci. U. S. A.* 96: 406-411 [Abstract] [Full Text]

- ▶ [Abstract of this Article](#)
- ▶ [Reprint \(PDF\) Version of this Article](#)
- ▶ Similar articles found in:
[PNAS Online](#)
[ISI Web of Science](#)
[PubMed](#)
- ▶ [PubMed Citation](#)
- ▶ This Article has been cited by:
- ▶ Search Medline for articles by:
[Sasaki, K. || Fukuda, M.](#)
- ▶ Search for citing articles in:
[ISI Web of Science \(18\)](#)
- ▶ Alert me when:
[new articles cite this article](#)
- ▶ [Download to Citation Manager](#)

[HOME](#) [HELP](#) [FEEDBACK](#) [SUBSCRIPTIONS](#) [ARCHIVE](#) [SEARCH](#)

Copyright (C) 1997 by the National Academy of Sciences